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1: AQ382839. RPCI11-139I6.TV R...[gi:4353862]   Links

```
LOCUS       AQ382839                626 bp    DNA     linear   GSS 21-MAY-1999
DEFINITION  RPCI11-139I6.TV RPCI-11 Homo sapiens genomic clone RPCI-11-139I6,
            DNA sequence.
ACCESSION   AQ382839
VERSION     AQ382839.1  GI:4353862
KEYWORDS    GSS.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 626)
  AUTHORS   Zhao,S., Adams,M.D., Nierman,W., Malek,J., de Jong,P. and Venter
            ,J.C.
  TITLE     Use of BAC End Sequences from Library RPCI-11 for Sequence-Ready
            Map Building
  JOURNAL   Unpublished (1997)
COMMENT     Other_GSSs: RPCI11-139I6.TJ
            Contact: Shaying Zhao, William Nierman, Mark Adams
            Department of Eukaryotic Genomics
            The Institute for Genomic Research
            9712 Medical Center Dr., Rockville, MD 20850
            Tel: 301 838 0200
            Fax: 301 838 0208
            Email: hbe@tigr.org
            Clones are derived from the human BAC library RPCI-11. For BAC
            library availability, please contact Pieter de Jong
            (pieter@dejong.med.buffalo.edu). Clones may be purchased from
            BACPAC Resources (http://bacpac.med.buffalo.edu/ordering) or from
            Research Genetics (info@resgen.com). BAC end search page:
            http://www.tigr.org/tdb/humgen/bac_end_search/bac_end_search.html
            Seq primer: T7
            Class: BAC ends.
FEATURES             Location/Qualifiers
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                     /clone="RPCI-11-139I6"
                     /clone_lib="RPCI-11"
                     /sex="Male"
                     /cell_type="Lymphocytes"
                     /note="Vector: pBACe3.6; Site_1: EcoRI; Site_2: EcoRI;
                     RPCI11 Human Male BAC Library"
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       61 aagcagtaaa acctgttcag acagcatcat tatattactt tcaggcaaac aagcgtggag
      121 gacctctaca ccctcaagtt aaaaaaggca ctgaggccgg gcgcggtggc tcacgcctgt
      181 aatcccaaca ctttgggagg ccaaggtggg tggatcacga ggtcaagaga tcgagaccat
      241 cctggccaac atggtgaaac cccgtctcta ctaaaaatac aaaaaaaata gccgggcatg
      301 gtggcgggtg cctgtagtgc cagctactcg ggaggctgag gcaggagaat ggtgtgaacc
      361 cgggaggcgg aacttgcagt gagccgagat cacgccactg cactccagcc tgggcaacag
      421 agtgagactg tctcaaaaaa aaaccaacca aacaaacaaa aaagcactga aagagacaga
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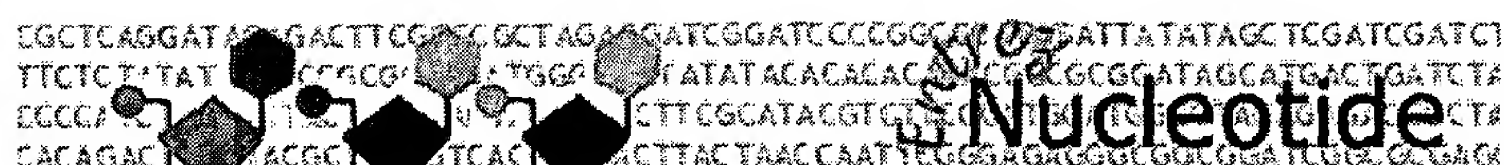
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481 ctccagttcc cggaaggctg acccagtcag cactaaaaac aggattaatg tatagctatt
541 aagagaatca tacagtggct ttattcttac tacttaaaaa aaggtgatgt gatggcagtg
601 atggtcaaca tcacacaggg aagacc
//
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Revised: July 5, 2002.

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☐ 1: AI692778. wd70d04.x1 NCI_CG...[gi:4970118]   Links

```
LOCUS       AI692778                565 bp    mRNA    linear   EST 17-DEC-1999
DEFINITION  wd70d04.x1 NCI_CGAP_Lu24 Homo sapiens cDNA clone IMAGE:2336935 3',
            mRNA sequence.
ACCESSION   AI692778
VERSION     AI692778.1  GI:4970118
KEYWORDS    EST.
SOURCE      Homo sapiens (human)
  ORGANISM  Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1  (bases 1 to 565)
  AUTHORS   NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
  TITLE     National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
            Tumor Gene Index
  JOURNAL   Unpublished (1997)
COMMENT     Contact: Robert Strausberg, Ph.D.
            Email: cgapbs-r@mail.nih.gov
            Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
            Emmert-Buck, M.D., Ph.D.
             cDNA Library Preparation: M. Bento Soares, Ph.D.
             cDNA Library Arrayed by: Greg Lennon, Ph.D.
             DNA Sequencing by: Washington University Genome Sequencing Center
             Clone distribution: NCI-CGAP clone distribution information can be
            found through the I.M.A.G.E. Consortium/LLNL at:
            www-bio.llnl.gov/bbrp/image/image.html
            Insert Length: 637    Std Error: 0.00
            Seq primer: -40UP from Gibco
            High quality sequence stop: 487.
FEATURES             Location/Qualifiers
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                     /db_xref="taxon:9606"
                     /clone="IMAGE:2336935"
                     /clone_lib="NCI_CGAP_Lu24"
                     /tissue_type="carcinoid"
                     /lab_host="DH10B"
                     /note="Organ: lung; Vector: pT7T3D-Pac (Pharmacia) with a
                     modified polylinker; Plasmid DNA from the normalized
                     library NCI_CGAP_Lu5 was prepared, and ss circles were
                     made in vitro. Following HAP purification, this DNA was
                     used as tracer in a subtractive hybridization reaction.
                     The driver was PCR-amplified cDNAs from a pool of 5,000
                     clones made from the same library (cloneIDs
                     1414920-1417991 and 1520904-1522439). Subtraction by Bento
                     Soares and M. Fatima Bonaldo. "
BASE COUNT      142 a    146 c    168 g    109 t
ORIGIN
        1 gcggccgcag cgctgaatgg ggtggaccga cgttccctgc agcgttcagc aaggctggct
       61 ctagaagtgc tggagagggc caagaggagg gcggtggact ggcatgccct ggagcgtccc
      121 aaaggctgca tggggggtcct tgcccgggag gcgcccccacc tagagaaaca gccggcagcc
      181 ggcccgcagc gcgttctccc gggagagaaa tattattcat ctgtgccaga ggaaggaggg
      241 gcaacccatg tctatcgtta tcacagaggc gagtcgaagc tgcacatgtg cttggacata
      301 gggaatggtc agagaaaaga cagaaaaaag acatcccttg gtcctggagg cagctatcaa
```

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        361 atatcagagc atgctccaga ggcatcccag cctgctgaga acatctctaa ggacctctac
        421 atagaagtat atccagggac ctattctgtc actgtgggct caaatgactt aaccaagaag
        481 actcatgtgg tagcagttga ttctggacaa agcgtggacc tggtcttccc tgtgtgatgt
        541 tgaccatcac tgccatcaca tcacc
//
```

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